

ID AQ743305 standard; DNA; GSS; 1141 BP.

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AC AQ743305;

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SV AQ743305.1

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DT 19-JUL-1999 (Rel. 60, Created)

DT 04-MAR-2000 (Rel. 63, Last updated, Version 2)

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DE HS_5387_B2_B02_SP6 RPCI-11 Human Male BAC Library Homo sapiens genomic
DE clone Plate=963 Col=4 Row=D, genomic survey sequence.

XX

KW GSS.

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OS Homo sapiens (human)

OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;

OC Eutheria; Primates; Catarrhini; Hominidae; Homo.

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RN [1]

RP 1-1141

RX MEDLINE; 99380589.

RA Mahairas G.G., Wallace J.C., Smith K., Swartzell S., Holzman T., Keller A.,

RA Shaker R., Furlong J., Young J., Zhao S., Adams M.D., Hood L.;

RT "Sequence-tagged connectors: A sequence approach to mapping and scanning
RT the human genome";

RL Proc. Natl. Acad. Sci. U.S.A. 96(17):9739-9744(1999).

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CC Contact: Mahairas GG, Wallace JC, Hood L

CC High Throughput Sequencing Center

CC University of Washington

CC 401 Queen Anne Avenue North, Seattle, WA 98109, USA

CC Tel: (206) 616-3618

CC Fax: (206) 616-3887

CC Email: jwallace@u.washington.edu

CC Clones are derived from the human BAC library RPCI-11. For BAC

CC library availability, please contact Pieter de Jong

CC (pieter@dejong.med.buffalo.edu). Clones may be purchased from

CC BACPAC Resources (http://bacpac.med.buffalo.edu/ordering_bac.htm)

CC or from Resear h Genetics (info@resgen.com). BAC end Web Server:

CC <http://www.htsc.washington.edu>

CC Plate: 963 row: D column: 4

CC Seq primer: SP6

CC Class: BAC ends

CC High quality sequence stop: 1141.

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FH Key Location/Qualifiers

FH

FT source

1..1141

FT

/db_xref="taxon:9606"

FT

/note="Vector: pBACe3.6; Genomic sequence of BAC ends"

FT

/sex="male"

FT

/organism="Homo sapiens"

FT

/clone="Plate=963 Col=4 Row=D"

FT

/clone_lib="RPCI-11 Human Male BAC Library"

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SQ Sequence 1141 BP; 760 A; 301 C; 16 G; 15 T; 49 other;

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attnnaaggg cggccaanaa anccctttan aaggaaaaan tacanggacc ncnnaaaana	120
nannaaaaaa aaaaaacaaa caaaaaatta aaaacaaaaa acaaaaaaaa ccacaaaaaa	180
cacacacaa naaaacacaa aaaacaaaaa acaaaaccac aacaacacaa caaaaaacca	240
aaacaaacaa aaaaaacaaa aaaaaaaaac aaaaaaaaac aaaaaaaaac ccanacacac	300
aacaacacaa caacaacacc aaaacacaca cacaacacnc accaaaaaaa caacaacaca	360
aaacacaaca aanacaaac aaaccacgac accaaacaac aacacacaca aaacacaaac	420
aaaaaacaaa aaaaaccaac aaacaaccaa aaaaaaaaac caaaaccaac aacacacaaa	480

caacaccaac	aacaacaaaa	caacaaaaca	aaaacaacaa	caaaccacaa	aaaaaacaaa	540
cacacaacaa	ncacaaaaaa	acaacaaaac	aaaacaacaa	aaaaaacaaa	acaaaaaaan	600
acaaaaacaa	caaacaacac	aaccacacaa	aacaacaaca	acaacaacaa	aacaacacca	660
ananaaacac	acaaaaaaca	aacaaaaaaa	acaaccaaca	aaaaacacaa	aacacaccca	720
aaaaaaaaacc	aaaaaaaaac	aanccaacaa	cccacaaaac	acacaaaaaa	aaaacaacca	780
cacacaacac	aacaaaccaa	aaaaaacccac	acacacaaac	caacaaaaca	aaacaaacaa	840
caaaccacac	caacaaaacc	acacaaacaa	ancaacaaca	caaacaacaa	accaacaann	900
cacaccacca	caaacacaca	caacaanaaa	accacaaaaa	cacacaaaac	aaaccacaaa	960
acaaaaccaa	ancaaccaac	aacaaaaaac	aacaacaaac	acaacaaaaa	aaacacaaaa	1020
acaaaaaana	ccaaaacaac	aaacaacaaa	caccacaaca	accaacaaaa	aacacaaanc	1080
aaacaaaaaa	acaaaaaaaa	aaaaaacaaa	aaaaacaaca	caacacaaaa	aacaaaaaaa	1140
a						1141

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